



```

Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match: 100.00%
DB: 10
Gaps: 0

US 09 856-070-25 (1-5) x US-09-834-975-688 (1-51)

QY 1 MetLeuArgLeuGln 5
bb 19 ATGCTACGCTTCAC 44

RESULT 2
US-09-974-300-4123
: Sequence 4123, Application US/09474300
: Patent No. US20020146721A1
: GENERAL INFORMATION:
: APPLICANT: Berk, Randy M.
: APPLICANT: Clausen, Ib Groth
: TITLE OF INVENTION: Methods For Monitoring Multiple Gene
: TITLE OF INVENTION: Expression
: FILE REFERENCE: 10085,530 US
: CURRENT APPLICATION NUMBER: US/09/974,300
: PRIOR FILING DATE: 2001-10-05
: PRIOR APPLICATION NUMBER: 09/680,598
: PRIOR FILING DATE: 2000-10-06
: PRIOR APPLICATION NUMBER: 60/279,526
: PRIOR FILING DATE: 2001-03-27
: NUMBER OF SEQ ID NOS: 8481
: SOFTWARE: FastSeq for Windows Version 4.0
: SEQ ID NO 4123
: LENGTH: 129
: TYPE: DNA
: ORGANISM: Bacillus licheniformis
US-09-974-300-4123

Alignment Scores:
Pred. No.: 108 Length: 129
Score: 23.00 Matches: 5
Percent Similarity: 100.00% Conservativeness: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
DB: 10 Gaps: 0

US 09 856-070-25 (1-5) x US-09-974-300-4123 (1-129)

QY 1 MetLeuArgLeuGln 5
bb 66 ATGTTACGACTTCAC 80

RESULT 3
US 09 864-761-20446
: Sequence 20446, Application US/09864761
: Patent No. US20020048763A1
: GENERAL INFORMATION:
: APPLICANT: Penn, Sharon G.
: APPLICANT: Rank, David R.
: APPLICANT: Hanzel, David K.
: APPLICANT: Chen, Wensheng
: TITLE OF INVENTION: HUMAN GENEOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
: TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
: FILE REFERENCE: Apmica-X-1
: CURRENT APPLICATION NUMBER: US/09/864,761
: PRIOR FILING DATE: 2001-05-23
: PRIOR APPLICATION NUMBER: US 60/180,312
: PRIOR FILING DATE: 2000-02-04
: PRIOR APPLICATION NUMBER: US 60/207,456
: PRIOR FILING DATE: 2000-05-26
: PRIOR APPLICATION NUMBER: US 60/643,466
: PRIOR FILING DATE: 2000-08-03
: PRIOR APPLICATION NUMBER: GB 2426346
: PRIOR FILING DATE: 2000-10-04
: PRIOR APPLICATION NUMBER: US 60/236,459
: PRIOR FILING DATE: 2000-09-27

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: PRIOR APPLICATION NUMBER: PCT/US01/00666
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCT/US01/00667
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCT/US01/00664
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCT/US01/00669
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCT/US01/00665
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCT/US01/00668
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCT/US01/00663
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCT/US01/00662
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCT/US01/00661
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCT/US01/00670
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: US 60/234,687
: PRIOR FILING DATE: 2000-09-21
: PRIOR APPLICATION NUMBER: US 09/608,408
: PRIOR FILING DATE: 2000-06-30
: PRIOR APPLICATION NUMBER: US 09/774,203
: PRIOR FILING DATE: 2001-01-29
: NUMBER OF SEQ ID NOS: 49117
: SOFTWARE: Anomax Sequence Listing Engine vers. 1.1
: SEQ ID NO 20446
: LENGTH: 139
: TYPE: DNA
: ORGANISM: Homo sapiens
: FEATURE:
: OTHER INFORMATION: MAP TO AF000409.1
: OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL - 18
: OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL - 2
: OTHER INFORMATION: NT HIT: g11423558, EVALUATE 1.20e+00
: OTHER INFORMATION: EST_HUMAN HIT: AA758580.1, EVALUATE 1.80e+00
: OTHER INFORMATION: SWISSPROT HIT: Q55029, EVALUATE 6.70e+00
US-09-864-761-20446

Alignment Scores:
Pred. No.: 117 Length: 139
Score: 23.00 Matches: 5
Percent Similarity: 100.00% Conservativeness: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
DB: 10 Gaps: 0

US-09-856-070-25 (1-5) x US-09-864-761-20446 (1-139)

QY 1 MetLeuArgLeuGln 5
bb 60 ATGCTTAGCTTCAC 74

RESULT 4
US-09-867-701-219
: Sequence 219, Application US/09867701
: Patent No. US2002013237A1
: GENERAL INFORMATION:
: APPLICANT: Aquile, Paul A.
: APPLICANT: Jones, Robert
: APPLICANT: Harlocker, Susan L.
: TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
: TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
: FILE REFERENCE: 210121.497
: CURRENT APPLICATION NUMBER: US/09/867,701
: PRIOR FILING DATE: 2001-05-29
: NUMBER OF SEQ ID NOS: 10912
: SOFTWARE: FastSeq for Windows Version 4.0
: SEQ ID NO 219
: LENGTH: 152
: TYPE: DNA

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: ORGANISM: Homo sapiens
US-09-867-701-219

Alignment Scores:
Pred. No.: 138 Length: 152
Score: 23.00 Matches: 5
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
Gaps: 0
DB: 10

US-09-856-070-25 (1-5) x US-09-867-701-219 (1-152)

QY 1 MetLeuArgLeuGln 5
|||||
DB 12 ATGCTGGGCTGGAG 26

RESULT 5
US-09-764-869-39
: Sequence 39, Application US/09/764869
: Patent No. US20020048763A1
: GENERAL INFORMATION:
: APPLICANT: Rosen et al.
: TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
: FILE REFERENCE: PC007
: CURRENT APPLICATION NUMBER: US/09/764,869
: CURRENT FILING DATE: 2001-01-17
: Prior application data removed - refer to PALM or file wrapper
: NUMBER OF SEQ ID NOS: 2442
: SOFTWARE: PatentIn Ver. 2.0
: SEQ ID NO 39
: LENGTH: 210
: TYPE: DNA
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: SITE
: LOCATION: (137)
: OTHER INFORMATION: n equals a,t,g, or c
US-09-764-869-39

Alignment Scores:
Pred. No.: 181 Length: 210
Score: 23.00 Matches: 5
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
Gaps: 0
DB: 10

US-09-856-070-25 (1-5) x US-09-764-869-39 (1-210)

QY 1 MetLeuArgLeuGln 5
|||||
DB 132 ATGCTGGGCTGGAG 146

RESULT 6
US-09-864-761-28662/c
: Sequence 28662, Application US/09/864761
: Patent No. US20020048763A1
: GENERAL INFORMATION:
: APPLICANT: Penn, Sharron G.
: APPLICANT: Rank, David R.
: APPLICANT: Hanzel, David K.
: APPLICANT: Chen, Wensheng
: TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
: FILE REFERENCE: Aconica-X-1
: CURRENT APPLICATION NUMBER: US/09/864,761
: CURRENT FILING DATE: 2001-05-23
: PRIOR APPLICATION NUMBER: US 6,071,800 A12
: PRIOR FILING DATE: 2000-02-04
: PRIOR APPLICATION NUMBER: US 6,020,456
: PRIOR FILING DATE: 2000-05-26
: PRIOR APPLICATION NUMBER: US 09/632,366

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: PRIOR FILING DATE: 2000-08-03
: PRIOR APPLICATION NUMBER: GB 24363.6
: PRIOR FILING DATE: 2000-10-04
: PRIOR APPLICATION NUMBER: US 60/236,359
: PRIOR FILING DATE: 2000-09-27
: PRIOR APPLICATION NUMBER: PCI/US01/00666
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCI/US01/00667
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCI/US01/00664
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCI/US01/00669
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCI/US01/00665
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCI/US01/00668
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCI/US01/00663
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCI/US01/00662
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCI/US01/00661
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: PCI/US01/00670
: PRIOR FILING DATE: 2001-01-30
: PRIOR APPLICATION NUMBER: US 60/234,687
: PRIOR FILING DATE: 2000-09-21
: PRIOR APPLICATION NUMBER: US 09/608,408
: PRIOR FILING DATE: 2000-06-30
: PRIOR APPLICATION NUMBER: US 09/774,203
: PRIOR FILING DATE: 2001-01-29
: NUMBER OF SEQ ID NOS: 49117
: SOFTWARE: Annonax Sequence Listing Engine vers. 1.1
: SEQ ID NO 28662
: LENGTH: 219
: TYPE: DNA
: ORGANISM: Homo sapiens
: FEATURE:
: OTHER INFORMATION: MAP TO AC010290.4
: OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1
: OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.8
: OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.85
: OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0.75
: OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 0.9
: OTHER INFORMATION: SWISSPROT HIT: Q91739, EVALUATE 1.90e-01
: OTHER INFORMATION: NT HIT: AB033057.1, EVALUATE 1.00e-120
US-09-864-761-28662

: Alignment Scores:
Pred. No.: 189 Length: 219
Score: 23.00 Matches: 5
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
Gaps: 0
DB: 10

US-09-856-070-25 (1-5) x US-09-864-761-28662 (1-219)

QY 1 MetLeuArgLeuGln 5
|||||
DB 150 ATGCTGGGCTGGAG 136

RESULT 7
US-09-864-761-30470
: Sequence 30470, Application US/09/864761
: Patent No. US20020048763A1
: GENERAL INFORMATION:
: APPLICANT: Penn, Sharron G.
: APPLICANT: Rank, David R.
: APPLICANT: Hanzel, David K.
: APPLICANT: Chen, Wensheng
: TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
: TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY

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FILE REFERENCE: Acomica-X-1  
CURRENT APPLICATION NUMBER: US/09/864,761  
CURRENT FILING DATE: 2001-05-23  
PRIOR APPLICATION NUMBER: US 60/180,312  
PRIOR FILING DATE: 2000-02-04  
PRIOR APPLICATION NUMBER: US 60/207,456  
PRIOR FILING DATE: 2000-05-26  
PRIOR APPLICATION NUMBER: US 09/642,356  
PRIOR FILING DATE: 2000-08-03  
PRIOR APPLICATION NUMBER: GR 24263.6  
PRIOR FILING DATE: 2000-10-04  
PRIOR APPLICATION NUMBER: US 60/236,359  
PRIOR FILING DATE: 2000-09-27  
PRIOR APPLICATION NUMBER: PCT/US01/00666  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: PCT/US01/00667  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: PCT/US01/00664  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: PCT/US01/00669  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: PCT/US01/00665  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: PCT/US01/00668  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: PCT/US01/00663  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: PCT/US01/00662  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: PCT/US01/00661  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: PCT/US01/00670  
PRIOR FILING DATE: 2001-01-30  
PRIOR APPLICATION NUMBER: US 60/234,687  
PRIOR FILING DATE: 2000-09-21  
PRIOR APPLICATION NUMBER: US 60/608,408  
PRIOR FILING DATE: 2000-06-30  
PRIOR APPLICATION NUMBER: US 60/774,203  
PRIOR FILING DATE: 2001-01-29  
NUMBER OF SEQ ID NOS: 49117  
SOFTWARE: Autobox Sequence Listing Engine vers. 1.1  
SEQ ID NO 30470  
LENGTH: 226  
TYPE: DNA  
ORGANISM: Homo sapiens  
FEATURE:  
OTHER INFORMATION: MAP TO APOBQ494.1  
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1  
OTHER INFORMATION: NT HIT: AJ012116.1, EVALUATE 1.40e-01  
OTHER INFORMATION: EST\_HUMAN HIT: R56395.1, EVALUATE 1.40e-01  
OTHER INFORMATION: SWISSPROT HIT: Q04490, EVALUATE 3.90e+00  
US-09-864,761-30470

Alignment Scores:  
Pred. No.: 195 Length: 226  
Score: 24.00 Matches: 5  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 100.00% Indels: 0  
DB: 10 Gaps: 0

US-09-856-070-25 (1-5) x US-09-864-761-30470 (1-226)

QY 1 MetLeuArgLeuGln 5  
|||||

DB 108 ATGTTGAGGTTGAG 122

RESULT 8

US-09-764-869-572/c  
Sequence 572, Application US/09764869  
Patent No. US20020061521A1  
GENERAL INFORMATION:  
APPLICANT: Rosen et al.

FILE REFERENCE: p007  
CURRENT APPLICATION NUMBER: US/09/764,869  
CURRENT FILING DATE: 2001-01-17  
Prior application data removed refer to PAM or file wrapper  
NUMBER OF SEQ ID NOS: 2442  
SOFTWARE: PatentIn Ver. 2.0  
SEQ ID NO 572  
LENGTH: 249  
TYPE: DNA  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: SITE  
LOCATION: (207)  
OTHER INFORMATION: n equals a,t,g, or c  
NAME/KEY: SITE  
LOCATION: (246)  
OTHER INFORMATION: n equals a,t,g, or c  
US-09-764-869-572

Alignment Scores:  
Pred. No.: 216 Length: 249  
Score: 24.00 Matches: 5  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 100.00% Indels: 0  
DB: 10 Gaps: 0

US-09-856-070-25 (1-5) x US-09-764-869-572 (1-249)

QY 1 MetLeuArgLeuGln 5

|||||

DB 58 AUGGUCAGATTACAG 44

RESULT 9

US-09-878-574-11486  
Sequence 11486, Application US/09878574  
Patent No. US20020110548A1

GENERAL INFORMATION:

APPLICANT: Hyrum, Joseph R.

APPLICANT: La Rosa, Thomas J.

APPLICANT: Thompson, Michael D.

FILE REFERENCE: 38-21(15401)B

FILE REFERENCE: 38-21(15401)B

CURRENT APPLICATION NUMBER: US/09/878,574

CURRENT FILING DATE: 2001-12-21

PRIOR APPLICATION NUMBER: 09/333,535

PRIOR FILING DATE: 1999-06-14

NUMBER OF SEQ ID NOS: 15775

SEQ ID NO 11486

LENGTH: 255

TYPE: DNA

ORGANISM: Glycine max

OTHER INFORMATION: Clone ID: 701064494H1

US-09-878-574-11486

Alignment Scores:  
Pred. No.: 222 Length: 255  
Score: 24.00 Matches: 5  
Percent Similarity: 100.00% Conservative: 0  
Best Local Similarity: 100.00% Mismatches: 0  
Query Match: 100.00% Indels: 0  
DB: 10 Gaps: 0

US-09-856-070-25 (1-5) x US-09-878-574-11486 (1-255)

QY 1 MetLeuArgLeuGln 5

|||||

DB 180 AUGGUCAGATTACAG 194

RESULT 10

US-10-046-935-1795

```

: Sequence 1795, Application US/10046935
: Patent No. US20020156011A1
: GENERAL INFORMATION:
: APPLICANT: Jiang, Yuqiu
: APPLICANT: Harlocker, Susan L.
: APPLICANT: Secrist, Heather
: APPLICANT: Wang, Aijun
: APPLICANT: Stolk, John A.
: TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
: TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
: FILE REFERENCE: 210121.527C1
: CURRENT APPLICATION NUMBER: US/10/046,935
: CURRENT FILING DATE: 2002-01-15
: NUMBER OF SEQ ID NOS: 2239
: SOFTWARE: FastSeq for Windows Version 4.0
: SEQ ID NO 1795
: LENGTH: 256
: TYPE: DNA
: ORGANISM: Homo sapiens
: FEATURE:
: NAME/KEY: misc_feature
: LOCATION: 228
: OTHER INFORMATION: n = A,T,C or G
US-10-046-935-1795

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```

Alignment Scores:
Pred. No.: 222 Length: 256
Score: 23.00 Matches: 5
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
DB: 9 Gaps: 0

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US-09-856-070-25 (1-5) x US-10-046-935-1795 (1-256)
```

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QY 1 MetLeuArqLeuGln 5
Db 14 ATGCTCAGGCTTCAG 28

RESULT 11
US-09-878-178-1795
: Sequence 1795, Application US/09878178
: Patent No. US20020177552A1
: GENERAL INFORMATION:
: APPLICANT: Jiang, Yuqiu
: APPLICANT: Harlocker, Susan L.
: APPLICANT: Secrist, Heather
: TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
: TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
: FILE REFERENCE: 210121.527
: CURRENT APPLICATION NUMBER: US/09/878,178
: CURRENT FILING DATE: 2001-06-08
: NUMBER OF SEQ ID NOS: 2237
: SOFTWARE: FastSeq for Windows Version 4.0
: SEQ ID NO 1795
: LENGTH: 256
: TYPE: DNA
: ORGANISM: Homo sapien
: FEATURE:
: NAME/KEY: misc_feature
: LOCATION: (1)...(256)
: OTHER INFORMATION: n = A,T,C or G
US-09-878-178-1795

```

```

Alignment Scores:
Pred. No.: 222 Length: 256
Score: 23.00 Matches: 5
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
DB: 9 Gaps: 0

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```
US-09-856-070-25 (1-5) x US-09-878-178-1795 (1-256)
```

```

QY 1 MetLeuArqLeuGln 5
Db 14 ATGCTCAGGCTTCAG 28

RESULT 12
US-10-046-935-289/c
: Sequence 289, Application US/10046935
: Patent No. US20020156011A1
: GENERAL INFORMATION:
: APPLICANT: Jiang, Yuqiu
: APPLICANT: Harlocker, Susan L.
: APPLICANT: Secrist, Heather
: APPLICANT: Wang, Aijun
: APPLICANT: Stolk, John A.
: TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
: TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
: FILE REFERENCE: 210121.527C1
: CURRENT APPLICATION NUMBER: US/10/046,935
: CURRENT FILING DATE: 2002-01-15
: NUMBER OF SEQ ID NOS: 2239
: SOFTWARE: FastSeq for Windows Version 4.0
: SEQ ID NO 289
: LENGTH: 257
: TYPE: DNA
: ORGANISM: Homo sapiens
US-10-046-935-289

```

```

Alignment Scores:
Pred. No.: 223 Length: 257
Score: 23.00 Matches: 5
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
DB: 9 Gaps: 0

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US-09-856-070-25 (1-5) x US-10-046-935-289 (1-257)
```

```

QY 1 MetLeuArqLeuGln 5
Db 243 ATGCTCAGGCTTCAG 229

RESULT 13
US-09-878-178-289/c
: Sequence 289, Application US/09878178
: Patent No. US20020177552A1
: GENERAL INFORMATION:
: APPLICANT: Jiang, Yuqiu
: APPLICANT: Harlocker, Susan L.
: APPLICANT: Secrist, Heather
: TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
: TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
: FILE REFERENCE: 210121.527
: CURRENT APPLICATION NUMBER: US/09/878,178
: CURRENT FILING DATE: 2001-06-08
: NUMBER OF SEQ ID NOS: 2237
: SOFTWARE: FastSeq for Windows Version 4.0
: SEQ ID NO 289
: LENGTH: 257
: TYPE: DNA
: ORGANISM: Homo sapien
US-09-878-178-289

```

```

Alignment Scores:
Pred. No.: 223 Length: 257
Score: 23.00 Matches: 5
Percent Similarity: 100.00% Conservative: 0
Best Local Similarity: 100.00% Mismatches: 0
Query Match: 100.00% Indels: 0
DB: 9 Gaps: 0

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```
US-09-856-070-25 (1-5) x US-09-878-178-289 (1-257)
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